

10/584207

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SEQUENCE LISTING

<110> Colgan, Terence J.
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Yang, Eric C.C.

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<150> US 60/630,990

<151> 2004-11-24

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 Cys Ala Leu Gly Pro Glu Val Ala Asn Val Ala Lys Phe Leu Cys Arg
 260 265 270
 Gln Ser Ser Gly Glu Asn Cys Asp Val Val Val Asn Thr Leu Gly Lys
 275 280 285
 Arg Ala Pro Ala Phe Glu Gly Arg Ile Leu Leu Asn Pro Gln Asp Lys
 290 295 300
 Asp Gly Ser Phe Ser Val Val Ile Thr Gly Leu Arg Lys Glu Asp Ala
 305 310 315 320
 Gly Arg Tyr Leu Cys Gly Ala His Ser Asp Gly Gln Leu Gln Glu Gly
 325 330 335
 Ser Pro Ile Gln Ala Trp Gln Leu Phe Val Asn Glu Glu Ser Thr Ile
 340 345 350
 Pro Arg Ser Pro Thr Val Val Lys Gly Val Ala Gly Ser Ser Val Ala

355					360					365					
Val	Leu	Cys	Pro	Tyr	Asn	Arg	Lys	Glu	Ser	Lys	Ser	Ile	Lys	Tyr	Trp
	370					375					380				
Cys	Leu	Trp	Glu	Gly	Ala	Gln	Asn	Gly	Arg	Cys	Pro	Leu	Leu	Val	Asp
385					390					395					400
Ser	Glu	Gly	Trp	Val	Lys	Ala	Gln	Tyr	Glu	Gly	Arg	Leu	Ser	Leu	Leu
				405					410					415	
Glu	Glu	Pro	Gly	Asn	Gly	Thr	Phe	Thr	Val	Ile	Leu	Asn	Gln	Leu	Thr
			420					425					430		
Ser	Arg	Asp	Ala	Gly	Phe	Tyr	Trp	Cys	Leu	Thr	Asn	Gly	Asp	Thr	Leu
		435					440					445			
Trp	Arg	Thr	Thr	Val	Glu	Ile	Lys	Ile	Ile	Glu	Gly	Glu	Pro	Asn	Leu
	450					455					460				
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Pro	Cys	His	Phe	Pro	Cys	Lys	Phe	Ser	Ser	Tyr	Glu	Lys	Tyr	Trp	Cys
				485					490					495	
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			500					505					510		
Pro	Ser	Lys	Ala	Phe	Val	Asn	Cys	Asp	Glu	Asn	Ser	Arg	Leu	Val	Ser
		515					520					525			
Leu	Thr	Leu	Asn	Leu	Val	Thr	Arg	Ala	Asp	Glu	Gly	Trp	Tyr	Trp	Cys
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Gly	Val	Lys	Gln	Gly	His	Phe	Tyr	Gly	Glu	Thr	Ala	Ala	Val	Tyr	Val
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Ala	Val	Glu	Glu	Arg	Lys	Ala	Ala	Gly	Ser	Arg	Asp	Val	Ser	Leu	Ala
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Lys	Ala	Asp	Ala	Ala	Pro	Asp	Glu	Lys	Val	Leu	Asp	Ser	Gly	Phe	Arg
			580					585					590		
Glu	Ile	Glu	Asn	Lys	Ala	Ile	Gln	Asp	Pro	Arg	Leu	Phe	Ala	Glu	Glu
		595					600					605			
Lys	Ala	Val	Ala	Asp	Thr	Arg	Asp	Gln	Ala	Asp	Gly	Ser	Arg	Ala	Ser
	610					615					620				

Val Asp Ser Gly Ser Ser Glu Glu Gln Gly Gly Ser Ser Arg Ala Leu
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 Val Ser Thr Leu Val Pro Leu Gly Leu Val Leu Ala Val Gly Ala Val
 645 650 655
 Ala Val Gly Val Ala Arg Ala Arg His Arg Lys Asn Val Asp Arg Val
 660 665 670
 Ser Ile Arg Ser Tyr Arg Thr Asp Ile Ser Met Ser Asp Phe Glu Asn
 675 680 685
 Ser Arg Glu Phe Gly Ala Asn Asp Asn Met Gly Ala Ser Ser Ile Thr
 690 695 700
 Gln Glu Thr Ser Leu Gly Gly Lys Glu Glu Phe Val Ala Thr Thr Glu
 705 710 715 720
 Ser Thr Thr Glu Thr Lys Glu Pro Lys Lys Ala Lys Arg Ser Ser Lys
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 Glu Glu Ala Glu Met Ala Tyr Lys Asp Phe Leu Leu Gln Ser Ser Thr
 740 745 750
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 <212> DNA
 <213> Homo sapiens

<300>
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 <309> 2004-10-27
 <313> (1)..(4266)

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 gctaaaacat tgcacaggag aagtcggcct gagtgggtgcg gcgctcggga cccaccagca 180
 atgctgctct tcgtgctcac ctgcctgctg gcggtcttcc cagccatctc cacgaagagt 240
 cccatatttg gtcccaggga ggtgaatagt gtggaaggta actcagtgtc catcacgtgc 300
 tactaccac ccacctctgt caaccggcac acccggaagt actgggtgccg gcaggagct 360
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 agggctaacc tcaccaactt cccggagaac ggcacatttg tggatgaacat tgcccagctg 480

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ggttatgtaa	atcccaacta	tacaggaaga	atacgccttg	atattcaggg	tactggccag	780
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caggctgggg	atgattccaa	tagtaataag	aagaatgctg	acctccaagt	gctaaagccc	900
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cctaggtttt tcctactgtc ctcagaggcg tgctggtccc ctcctcagtg acatcaaagc	2640
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attgcc	4266

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<213> Homo sapiens

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<309> 2004-10-25
<313> (1)..(187)

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Ala Val Asp Glu Leu Gly Lys Val Leu Thr Pro Thr Gln Val Lys Asn
35 40 45

Arg Pro Thr Ser Ile Ser Trp Asp Gly Leu Asp Ser Gly Lys Leu Tyr
50 55 60

Thr Leu Val Leu Thr Asp Pro Asp Ala Pro Ser Arg Lys Asp Pro Lys
65 70 75 80

Tyr Arg Glu Trp His His Phe Leu Val Val Asn Met Lys Gly Asn Asp
85 90 95

Ile Ser Ser Gly Thr Val Leu Ser Asp Tyr Val Gly Ser Gly Pro Pro
100 105 110

Lys Gly Thr Gly Leu His Arg Tyr Val Trp Leu Val Tyr Glu Gln Asp
115 120 125

Arg Pro Leu Lys Cys Asp Glu Pro Ile Leu Ser Asn Arg Ser Gly Asp
130 135 140

His Arg Gly Lys Phe Lys Val Ala Ser Phe Arg Lys Lys Tyr Glu Leu
145 150 155 160

Arg Ala Pro Val Ala Gly Thr Cys Tyr Gln Ala Glu Trp Asp Asp Tyr
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Val Pro Lys Leu Tyr Glu Gln Leu Ser Gly Lys
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<212> DNA
<213> Homo sapiens

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<308> NM_002567
<309> 2004-10-27
<313> (1)..(1507)

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gctgcacgct ctgcttgggc tcgccatgcc ggtggacctc agcaagtggg ccggggccctt 180
gagcctgcaa gaagtggacg agcagccgca gcacccgctg catgtcacct acgccggggc 240
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catttcgtgg gatggtcttg attcaggga gctctacacc ttggtcctga cagaccgga 360
tgctcccagc aggaaggatc ccaaatacag agaatggcat catttcctgg tggtaacat 420
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aaaaaaaaaa aaaaagattg gttgcctctg cttttgtgat cctgagtcca gaatggtaca 1140
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aaaaaaa 1507

<210> 13
<211> 249
<212> PRT
<213> Homo sapiens

<300>
<308> P39687

<309> 2004-06-15
<313> (1)..(249)

<400> 13

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			20					25					30		
Lys	Leu	Glu	Gly	Leu	Thr	Asp	Glu	Phe	Glu	Glu	Leu	Glu	Phe	Leu	Ser
		35					40					45			
Thr	Ile	Asn	Val	Gly	Leu	Thr	Ser	Ile	Ala	Asn	Leu	Pro	Lys	Leu	Asn
	50					55					60				
Lys	Leu	Lys	Lys	Leu	Glu	Leu	Ser	Asp	Asn	Arg	Val	Ser	Gly	Gly	Leu
65					70					75					80
Glu	Val	Leu	Ala	Glu	Lys	Cys	Pro	Asn	Leu	Thr	His	Leu	Asn	Leu	Ser
				85					90					95	
Gly	Asn	Lys	Ile	Lys	Asp	Leu	Ser	Thr	Ile	Glu	Pro	Leu	Lys	Lys	Leu
			100					105					110		
Glu	Asn	Leu	Lys	Ser	Leu	Asp	Leu	Phe	Asn	Cys	Glu	Val	Thr	Asn	Leu
		115					120					125			
Asn	Asp	Tyr	Arg	Glu	Asn	Val	Phe	Lys	Leu	Leu	Pro	Gln	Leu	Thr	Tyr
	130					135					140				
Leu	Asp	Gly	Tyr	Asp	Arg	Asp	Asp	Lys	Glu	Ala	Pro	Asp	Ser	Asp	Ala
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Glu	Gly	Tyr	Val	Glu	Gly	Leu	Asp	Asp	Glu	Glu	Glu	Asp	Glu	Asp	Glu
				165					170					175	
Glu	Glu	Tyr	Asp	Glu	Asp	Ala	Gln	Val	Val	Glu	Asp	Glu	Glu	Asp	Glu
			180					185					190		
Asp	Glu	Glu	Glu	Glu	Gly	Glu	Glu	Glu	Asp	Val	Ser	Gly	Glu	Glu	Glu
		195					200					205			
Glu	Asp	Glu	Glu	Gly	Tyr	Asn	Asp	Gly	Glu	Val	Asp	Asp	Glu	Glu	Asp
	210					215					220				
Glu	Glu	Glu	Leu	Gly	Glu	Glu	Glu	Arg	Gly	Gln	Lys	Arg	Lys	Arg	Glu
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<210> 14
<211> 1136
<212> DNA
<213> Homo sapiens

<300>
<308> NM_006305
<309> 2004-10-27
<313> (1)..(1136)

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gagagcgcga gagatggaga tgggcagacg gattcattta gagctgcgga acaggacgcc 180
ctctgatgtg aaagaacttg tcctggacaa cagtcgggtcg aatgaaggca aactcgaagg 240
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aatcgcaaac ttaccaaagt taaacaaact taagaagctt gaactaagcg ataacagagt 360
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gaaaaattcc tattgtgatt tgactgtttt taccatatac ccctctcccc cccccctcca 960
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<211> 643
<212> PRT
<213> Homo sapiens

<300>
<308> P17066
<309> 2003-09-15
<313> (1)..(643)

<400> 15

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 35 40 45
 Thr Glu Arg Leu Val Gly Asp Ala Ala Lys Ser Gln Ala Ala Leu Asn
 50 55 60
 Pro His Asn Thr Val Phe Asp Ala Lys Arg Leu Ile Gly Arg Lys Phe
 65 70 75 80
 Ala Asp Thr Thr Val Gln Ser Asp Met Lys His Trp Pro Phe Arg Val
 85 90 95
 Val Ser Glu Gly Gly Lys Pro Lys Val Arg Val Cys Tyr Arg Gly Glu
 100 105 110
 Asp Lys Thr Phe Tyr Pro Glu Glu Ile Ser Ser Met Val Leu Ser Lys
 115 120 125
 Met Lys Glu Thr Ala Glu Ala Tyr Leu Gly Gln Pro Val Lys His Ala
 130 135 140
 Val Ile Thr Val Pro Ala Tyr Phe Asn Asp Ser Gln Arg Gln Ala Thr
 145 150 155 160
 Lys Asp Ala Gly Ala Ile Ala Gly Leu Asn Val Leu Arg Ile Ile Asn
 165 170 175
 Glu Pro Thr Ala Ala Ala Ile Ala Tyr Gly Leu Asp Arg Arg Gly Ala
 180 185 190
 Gly Glu Arg Asn Val Leu Ile Phe Asp Leu Gly Gly Gly Thr Phe Asp
 195 200 205
 Val Ser Val Leu Ser Ile Asp Ala Gly Val Phe Glu Val Lys Ala Thr
 210 215 220
 Ala Gly Asp Thr His Leu Gly Gly Glu Asp Phe Asp Asn Arg Leu Val
 225 230 235 240
 Asn His Phe Met Glu Glu Phe Arg Arg Lys His Gly Lys Asp Leu Ser
 245 250 255

Gly Asn Lys Arg Ala Leu Arg Arg Leu Arg Thr Ala Cys Glu Arg Ala
 260 265 270
 Lys Arg Thr Leu Ser Ser Ser Thr Gln Ala Thr Leu Glu Ile Asp Ser
 275 280 285
 Leu Phe Glu Gly Val Asp Phe Tyr Thr Ser Ile Thr Arg Ala Arg Phe
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 Glu Glu Leu Cys Ser Asp Leu Phe Arg Ser Thr Leu Glu Pro Val Glu
 305 310 315 320
 Lys Ala Leu Arg Asp Ala Lys Leu Asp Lys Ala Gln Ile His Asp Val
 325 330 335
 Val Leu Val Gly Gly Ser Thr Arg Ile Pro Lys Val Gln Lys Leu Leu
 340 345 350
 Gln Asp Phe Phe Asn Gly Lys Glu Leu Asn Lys Ser Ile Asn Pro Asp
 355 360 365
 Glu Ala Val Ala Tyr Gly Ala Ala Val Gln Ala Ala Val Leu Met Gly
 370 375 380
 Asp Lys Cys Glu Lys Val Gln Asp Leu Leu Leu Asp Val Ala Pro
 385 390 395 400
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 405 410 415
 Gln Arg Asn Ala Thr Ile Pro Thr Lys Gln Thr Gln Thr Phe Thr Thr
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 Tyr Ser Asp Asn Gln Pro Gly Val Phe Ile Gln Val Tyr Glu Gly Glu
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 465 470 475 480
 Asp Ile Asp Ala Asn Gly Ile Leu Ser Val Thr Ala Thr Asp Arg Ser
 485 490 495
 Thr Gly Lys Ala Asn Lys Ile Thr Ile Thr Asn Asp Lys Gly Arg Leu
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 Ser Lys Glu Glu Val Glu Arg Met Val His Glu Ala Glu Gln Tyr Lys

515	520	525
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Arg Asp Lys Ile Pro Glu Glu Asp Arg Arg Lys Met Gln Asp Lys Cys 565 570 575		
Arg Glu Val Leu Ala Trp Leu Glu His Asn Gln Leu Ala Glu Lys Glu 580 585 590		
Glu Tyr Glu His Gln Lys Arg Glu Leu Glu Gln Ile Cys Arg Pro Ile 595 600 605		
Phe Ser Arg Leu Tyr Gly Gly Pro Gly Val Pro Gly Gly Ser Ser Cys 610 615 620		
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Glu Val Asp		

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 <213> Homo sapiens

<300>
 <308> NM_002155
 <309> 2004-10-28
 <313> (1)..(2664)

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 <213> Homo sapiens

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 <308> x51757
 <309> 1998-11-13
 <313> (1)..(2492)

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 <212> PRT
 <213> Homo sapiens

<300>
 <308> P14174
 <309> 2005-01-25
 <313> (1)..(115)

<400> 18

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Lys	Pro	Pro	Gln	Tyr	Ile	Ala	Val	His	Val	Val	Pro	Asp	Gln	Leu	Met
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Ala	Phe	Gly	Gly	Ser	Ser	Glu	Pro	Cys	Ala	Leu	Cys	Ser	Leu	His	Ser
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Ile Gly Lys Ile Gly Gly Ala Gln Asn Arg Ser Tyr Ser Lys Leu Leu
65 70 75 80

Cys Gly Leu Leu Ala Glu Arg Leu Arg Ile Ser Pro Asp Arg Val Tyr
85 90 95

Ile Asn Tyr Tyr Asp Met Asn Ala Ala Asn Val Gly Trp Asn Asn Ser
100 105 110

Thr Phe Ala
115

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<213> Homo sapiens

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<309> 2004-10-26
<313> (1)..(561)

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<212> DNA
<213> Homo sapiens

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<309> 1994-09-29
<313> (1)..(2167)

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<213> Homo sapiens

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<309> 2004-10-25
<313> (1)..(105)

<400> 21

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Thr Leu Ser Lys Thr Glu Phe Leu Ser Phe Met Asn Thr Glu Leu Ala
35 40 45

Ala Phe Thr Lys Asn Gln Lys Asp Pro Gly Val Leu Asp Arg Met Met
50 55 60

Lys Lys Leu Asp Thr Asn Ser Asp Gly Gln Leu Asp Phe Ser Glu Phe
65 70 75 80

Leu Asn Leu Ile Gly Gly Leu Ala Met Ala Cys His Asp Ser Phe Leu
85 90 95

Lys Ala Val Pro Ser Gln Lys Arg Thr
100 105

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<212> DNA
<213> Homo sapiens

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<309> 2004-10-26
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<400> 22

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 <213> Homo sapiens

<300>
 <308> P00938
 <309> 2003-09-15
 <313> (1)..(249)

<400> 23

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35 40 45

Ile Asp Phe Ala Arg Gln Lys Leu Asp Pro Lys Ile Ala Val Ala Ala
50 55 60

Gln Asn Cys Tyr Lys Val Thr Asn Gly Ala Phe Thr Gly Glu Ile Ser
65 70 75 80

Pro Gly Met Ile Lys Asp Cys Gly Ala Thr Trp Val Val Leu Gly His
85 90 95

Ser Glu Arg Arg His Val Phe Gly Glu Ser Asp Glu Leu Ile Gly Gln
100 105 110

Lys Val Ala His Ala Leu Ala Glu Gly Leu Gly Val Ile Ala Cys Ile
115 120 125

Gly Glu Lys Leu Asp Glu Arg Glu Ala Gly Ile Thr Glu Lys Val Val
130 135 140

Phe Glu Gln Thr Lys Val Ile Ala Asp Asn Val Lys Asp Trp Ser Lys
145 150 155 160

Val Val Leu Ala Tyr Glu Pro Val Trp Ala Ile Gly Thr Gly Lys Thr

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Ala Thr Pro	Gln Gln Ala Gln Glu Val	His Glu Lys Leu Arg	Gly Trp		
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Leu Lys Ser	Asn Val Ser Asp Ala Val	Ala Gln Ser Thr	Arg Ile Ile		
	195	200	205		
Tyr Gly Gly	Ser Val Thr Gly Ala Thr	Cys Lys Glu Leu	Ala Ser Gln		
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	245				

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 <212> DNA
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<300>
 <308> x69723
 <309> 1997-03-24
 <313> (1)..(5005)

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Ala Asn Lys Arg His Gly Ser Trp Lys Ile Gln Leu Asn Ala Thr Ser
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Val Thr His Lys Pro Asn Ala Ile Gln Met Ala Leu Ser Val Cys Glu
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Asp Leu Ile Ser Ser Gln Val Tyr Ala Ile Leu Val Ser His Pro Pro
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Thr Pro Asn Asp His Phe Thr Pro Thr Pro Val Ser Tyr Thr Ala Gly
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Phe Tyr Arg Ile Pro Val Leu Gly Leu Thr Thr Arg Met Ser Ile Tyr
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Ser Asp Lys Ser Ile His Leu Ser Phe Leu Arg Thr Val Pro Pro Tyr
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Ser His Gln Ser Ser Val Trp Phe Glu Met Met Arg Val Tyr Ser Trp
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Asn His Ile Ile Leu Leu Val Ser Asp Asp His Glu Gly Arg Ala Ala
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Gln Lys Arg Leu Glu Thr Leu Leu Glu Glu Arg Glu Ser Lys Ala Glu
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Lys Val Leu Gln Phe Asp Pro Gly Thr Lys Asn Val Thr Ala Leu Leu
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Met Glu Ala Lys Glu Leu Glu Ala Arg Val Ile Ile Leu Ser Ala Ser
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Lys Ile Thr Pro Asn Leu Ala Glu Phe Ala Phe Ser Leu Tyr Arg Gln
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Leu Ala His Gln Ser Asn Ser Thr Asn Ile Phe Phe Ser Pro Val Ser
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Ile Ala Thr Ala Phe Ala Met Leu Ser Leu Gly Thr Lys Ala Asp Thr
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His Asp Glu Ile Leu Glu Gly Leu Asn Phe Asn Leu Thr Glu Ile Pro
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Glu Ala Gln Ile His Glu Gly Phe Gln Glu Leu Leu Arg Thr Leu Asn
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Ser Glu Gly Leu Lys Leu Val Asp Lys Phe Leu Glu Asp Val Lys Lys
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Leu Tyr His Ser Glu Ala Phe Thr Val Asn Phe Gly Asp Thr Glu Glu
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Ala Lys Lys Gln Ile Asn Asp Tyr Val Glu Lys Gly Thr Gln Gly Lys
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Val Asn Tyr Ile Phe Phe Lys Gly Lys Trp Glu Arg Pro Phe Glu Val
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Lys Asp Thr Glu Glu Glu Asp Phe His Val Asp Gln Val Thr Thr Val
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Lys Lys Leu Ser Ser Trp Val Leu Leu Met Lys Tyr Leu Gly Asn Ala
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Thr Ala Ile Phe Phe Leu Pro Asp Glu Gly Lys Leu Gln His Leu Glu
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Asn Glu Leu Thr His Asp Ile Ile Thr Lys Phe Leu Glu Asn Glu Asp
290 295 300

Arg Arg Ser Ala Ser Leu His Leu Pro Lys Leu Ser Ile Thr Gly Thr
305 310 315 320

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Cys Thr Ile Gly Pro Ala Ser Arg Ser Val Glu Thr Leu Lys Glu Met
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Ile Lys Ser Gly Met Asn Val Ala Arg Leu Asn Phe Ser His Gly Thr
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His Glu Tyr His Ala Glu Thr Ile Lys Asn Val Arg Thr Ala Thr Glu
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Thr Ala Glu Val Glu Leu Lys Lys Gly Ala Thr Leu Lys Ile Thr Leu
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Tyr Lys Asn Ile Cys Lys Val Val Glu Val Gly Ser Lys Ile Tyr Val
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Glu Thr Leu Gln Leu Asp Ala Ala Gln His Ser Pro Ala Ser Pro Cys
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 115 120 125
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 355 360 365
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 515 520 525

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595 600 605

His Ser Leu Cys Ile Gln Glu Val Phe Pro Glu Asp Thr Gly Thr Tyr
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645 650 655

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660 665 670

Cys Ala Ile Ala Gly Asp Pro Phe Pro Thr Val His Trp Leu Arg Asp
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Glu Asp Val Phe Thr Leu Val Leu Lys Lys Val Gln Pro Trp His Ala
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725 730 735

Gln Val Ser Leu Met Leu Gln Asn Ser Ser Ala Arg Ala Leu Pro Arg
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755 760 765

Ala Asp Gly Gly Gly Ser Asp Arg Tyr Gly Ser Leu Arg Pro Gly Trp
770 775 780

Pro Ala Arg Gly Gln Gly Trp Leu Glu Glu Glu Asp Gly Glu Asp Val

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His 1040	Val Ala Glu Gly Lys	Lys 1045 Leu Leu Leu Gln	Cys 1050 Gln Val Ser

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Lys Leu Lys Gly Thr Glu Asp Glu Leu Asp Lys Tyr Ser Glu Ala Leu
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Ala Glu Ala Asp Val Ala Ser Leu Asn Arg Arg Ile Gln Leu Val Glu
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Glu Glu Leu Asp Arg Ala Gln Glu Arg Leu Ala Thr Ala Leu Gln Lys



100 105 110

Leu Glu Glu Ala Glu Lys Ala Ala Asp Glu Ser Glu Arg Gly Met Lys
115 120 125

Val Ile Glu Ser Arg Ala Gln Lys Asp Glu Glu Lys Met Glu Ile Gln
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Ile Lys Val Leu Ser Asp Lys Leu Lys Glu Ala Glu Thr Arg Ala Glu
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